



## SEQUENCE LISTING

<110> Ross, Theodora  
Mizukami, Ikuko

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<141> 2004-01-29

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 <222> (1267) .. (1268)  
  
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Met	Phe	Asp	Tyr	Leu	Glu	Cys	Glu	Leu	Asn	Leu	Phe	Gln	Thr	Val	Phe
1				5					10				15		

  

Asn	Ser	Leu	Asp	Met	Ser	Arg	Ser	Val	Ser	Val	Thr	Ala	Ala	Gly	Gln
				20				25				30			

  

Cys	Arg	Leu	Ala	Pro	Leu	Ile	Gln	Val	Ile	Leu	Asp	Cys	Ser	His	Leu
						40					45				

  

Tyr	Asp	Tyr	Thr	Val	Lys	Leu	Leu	Phe	Lys	Leu	His	Ser	Cys	Leu	Pro
				50			55			60					

  

Ala	Asp	Thr	Leu	Gln	Gly	His	Arg	Asp	Arg	Phe	Met	Glu	Gln	Phe	Thr
65					70				75				80		

  

Lys	Leu	Lys	Asp	Leu	Phe	Tyr	Arg	Ser	Ser	Asn	Leu	Gln	Tyr	Phe	Lys
					85			90				95			

  

Arg	Leu	Ile	Gln	Ile	Pro	Gln	Leu	Pro	Glu	Asn	Pro	Pro	Asn	Phe	Leu
					100			105				110			

  

Arg	Ala	Ser	Ala	Leu	Ser	Glu	His	Ile	Ser	Pro	Val	Val	Val	Ile	Pro
						115		120				125			

  

Ala	Glu	Ala	Ser	Ser	Pro	Asp	Ser	Glu	Pro	Val	Leu	Glu	Lys	Asp	Asp
					130			135			140				

  

Leu	Met	Asp	Met	Asp	Ala	Ser	Gln	Gln	Asn	Leu	Phe	Asp	Asn	Lys	Phe
						145		150			155			160	

  

Asp	Asp	Ile	Phe	Gly	Ser	Ser	Phe	Ser	Ser	Asp	Pro	Phe	Asn	Phe	Asn
					165			170				175			

Ser Gln Asn Gly Val Asn Lys Asp' Glu Lys Asp His Leu Ile Glu Arg  
180 185 190

Leu Tyr Arg Glu Ile Ser Gly Leu Lys Ala Gln Leu Glu Asn Met Lys  
195 200 205

Thr Glu Ser Gln Arg Val Val Leu Gln Leu Lys Gly His Val Ser Glu  
210 215 220

Leu Glu Ala Asp Leu Ala Glu Gln Gln His Leu Arg Gln Gln Ala Ala  
225 230 235 240

Asp Asp Cys Glu Phe Leu Arg Ala Glu Leu Asp Glu Leu Arg Arg Gln  
245 250 255

Arg Glu Asp Thr Glu Lys Ala Gln Arg Ser Leu Ser Glu Ile Glu Arg  
260 265 270

Lys Ala Gln Ala Asn Glu Gln Arg Tyr Ser Lys Leu Lys Glu Lys Tyr  
275 280 285

Ser Glu Leu Val Gln Asn His Ala Asp Leu Leu Arg Lys Asn Ala Glu  
290 295 300

Val Thr Lys Gln Val Ser Met Ala Arg Gln Ala Gln Val Asp Leu Glu  
305 310 315 320

Arg Glu Lys Lys Glu Leu Glu Asp Ser Leu Glu Arg Ile Ser Asp Gln  
325 330 335

Gly Gln Arg Lys Thr Gln Glu Gln Leu Glu Val Leu Glu Ser Leu Lys  
340 345 350

Gln Glu Leu Ala Thr Ser Gln Arg Glu Leu Gln Val Leu Gln Gly Ser  
355 360 365

Leu Glu Thr Ser Ala Gln Ser Glu Ala Asn Trp Ala Ala Glu Phe Ala  
370 375 380

Glu Leu Glu Lys Glu Arg Asp Ser Leu Val Ser Gly Ala Ala His Arg  
385 390 395 400

Glu Glu Glu Leu Ser Ala Leu Arg Lys Glu Leu Gln Asp Thr Gln Leu  
405 410 415

Lys Leu Ala Ser Thr Glu Glu Ser Met Cys Gln Leu Ala Lys Asp Gln  
420 425 430

Arg Lys Met Leu Leu Val Gly Ser Arg Lys Ala Ala Glu Gln Val Ile  
435 440 445

Gln Asp Ala Leu Asn Gln Leu Glu Glu Pro Pro Leu Ile Ser Cys Ala  
450 455 460

Gly Ser Ala Asp His Leu Leu Ser Thr Val Thr Ser Ile Ser Ser Cys  
465 470 475 480

Ile Glu Gln Leu Glu Lys Ser Trp Ser Gln Tyr Leu Ala Cys Pro Glu  
485 490 495

Asp Ile Ser Gly Leu Leu His Ser Ile Thr Leu Leu Ala His Leu Thr  
500 505 510

Ser Asp Ala Ile Ala His Gly Ala Thr Thr Cys Leu Arg Ala Pro Pro  
515 520 525

Glu Pro Ala Asp Ser Leu Thr Glu Ala Cys Lys Gln Tyr Gly Arg Glu  
530 535 540

Thr Leu Ala Tyr Leu Ala Ser Leu Glu Glu Gly Ser Leu Glu Asn  
545 550 555 560

Ala Asp Ser Thr Ala Met Arg Asn Cys Leu Ser Lys Ile Lys Ala Ile  
565 570 575

Gly Glu Glu Leu Leu Pro Arg Gly Leu Asp Ile Lys Gln Glu Glu Leu  
580 585 590

Gly Asp Leu Val Asp Lys Glu Met Ala Ala Thr Ser Ala Ala Ile Glu  
595 600 605

Thr Ala Thr Ala Arg Ile Glu Glu Met Leu Ser Lys Ser Arg Ala Gly  
610 615 620

Asp Thr Gly Val Lys Leu Glu Val Asn Glu Arg Ile Leu Gly Cys Cys  
625 630 635 640

Thr Ser Leu Met Gln Ala Ile Gln Val Leu Ile Val Ala Ser Lys Asp  
645 650 655

Leu Gln Arg Glu Ile Val Glu Ser Gly Arg Gly Thr Ala Ser Pro Lys  
660 665 670

Glu Phe Tyr Ala Lys Asn Ser Arg' Trp Thr Glu Gly Leu Ile Ser Ala  
675 680 685

Ser Lys Ala Val Gly Trp Gly Ala Thr Val Met Val Asp Ala Ala Asp  
690 695 700

Leu Val Val Gln Gly Arg Gly Lys Phe Glu Glu Leu Met Val Cys Ser  
705 710 715 720

His Glu Ile Ala Ala Ser Thr Ala Gln Leu Val Ala Ala Ser Lys Val  
725 730 735

Lys Ala Asp Lys Asp Ser Pro Asn Leu Ala Gln Leu Gln Gln Ala Ser  
740 745 750

Arg Gly Val Asn Gln Ala Thr Ala Gly Val Val Ala Ser Thr Ile Ser  
755 760 765

Gly Lys Ser Gln Ile Glu Glu Thr Asp Asn Met Asp Phe Ser Ser Met  
770 775 780

Thr Leu Thr Gln Ile Lys Arg Gln Glu Met Asp Ser Gln Val Arg Val  
785 790 795 800

Leu Glu Leu Glu Asn Glu Leu Gln Lys Glu Arg Gln Lys Leu Gly Glu  
805 810 815

Leu Arg Lys Lys His Tyr Glu Leu Ala Gly Val Ala Glu Gly Trp Glu  
820 825 830

Glu Gly Thr Glu Ala Ser Pro Pro Thr Leu Gln Glu Val Val Thr Glu  
835 840 845

Lys Glu Ser Gln Thr Asn Thr Pro Tyr Val Ser Val Asn Pro Cys Tyr  
850 855 860

Leu Ser Arg Val Cys Tyr Phe Pro Ser His Arg Pro Asn Pro Trp Ser  
865 870 875 880

Pro Arg Gly Ser His Thr Thr Ala Ile Thr Gln Cys Arg Gly His Ala  
885 890 895

His Phe Gln Arg Leu Pro Pro Arg His Pro Phe Cys Leu Asp Pro Trp  
900 905 910

Ile Ser Thr Ala Ser Tyr Gly Gly Trp Leu Gly Phe Leu Val Leu Phe  
915 920 925

Phe Phe Phe Lys Phe His Ser His Ser Gln Leu Ser Gln Arg Ala His  
930 935 940

Pro Trp Gly Val Ser Arg Ala Pro Gln Leu Trp Leu Gln Arg Trp Cys  
945 950 955 960

Cys Pro Gly Leu Ser Val Leu His Leu Arg Leu His Thr Asp Gln Val  
965 970 975

Leu Ala His Pro Val His Ala Pro Gly Ser Gly Gly Ala Ala Glu Gln  
980 985 990

Leu Ser Ser Lys Ser Arg Arg Arg Val Ser Ala Phe Pro Ser Ser Ile  
995 1000 1005

Pro Ala Glu Ser Leu Cys Pro Pro Leu Gln Gly Arg Arg Gln Gln  
1010 1015 1020

Lys Glu Gly Gln Glu Gly Ser His Ser Pro Val Pro Val Thr Arg  
1025 1030 1035

Leu Lys Asn Leu Ile Thr Cys Leu Asn Gly Ala Gly Glu Ile Asn  
1040 1045 1050

Asn Thr Thr Ser Leu Pro Glu Thr Val Arg Glu Trp Ser Leu Ser  
1055 1060 1065

Ser Gly Pro Ser Pro Leu Ala Gln Arg Arg Ser Val Gly Val Ile  
1070 1075 1080

Pro Asn Ser Phe Leu Gln Thr Ser Ala Leu Ala Ser Ser Ile Gly  
1085 1090 1095

Arg Ser Phe His Leu Leu Arg Asn Gln Thr Arg Lys Ile Arg Cys  
1100 1105 1110

Asn Cys Ser His Gln Gly Arg Thr Leu Tyr Leu Val Cys Tyr Pro  
1115 1120 1125

Tyr Leu Leu Leu Thr Ser Leu Lys Gln Gln Gln Pro Thr Lys Arg  
1130 1135 1140

Cys Leu Glu Gln Ser Glu Leu Gln Val Leu Gln Ser Ser Ser Phe  
1145 1150 1155

Cys	Pro	Ala	Thr	Ser	Ala	Phe	Lys	Asn	Gln	Lys	Lys	Gly	Gln	Gly
1160						1165					1170			
Ala	Gly	Leu	Leu	Leu	Thr	Trp	Ile	Pro	Lys	Gln	Gly	Asp	His	Leu
1175						1180					1185			
Glu	Leu	Leu	Gly	Gln	Arg	Lys	Glu	Arg	Thr	Glu	Pro	Ala	Ala	Pro
1190						1195					1200			
Thr	Pro	Phe	Ser	His	Met	Pro	Gln	Ala	Leu	Ala	Ala	Leu	Trp	Thr
1205						1210					1215			
Gly	Gly	Gln	Arg	Ala	His	Glu	Gln	Leu	Ala	Arg	Asp	Gly	Gln	Pro
1220						1225					1230			
Asn	Ser	Thr	Phe	Pro	Leu	Leu	Asp	Gly	Pro	Gln	His	Leu	Ser	Asp
1235						1240					1245			
Leu	Leu	Ile	Leu	Gly	Lys	Gln	Arg	Leu	Pro	Ser	Leu	Ser	Ile	Ala
1250						1255					1260			
Thr	His	Trp	Trp	Pro	Ser	Ser	Thr	Ser	Glu	Phe	Leu	Gln	Pro	Gly
1265						1270					1275			
Arg	Pro	Leu	Glu	His										
1280														
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<212>	DNA													
<213>	Homo sapiens													
<400>	5													
	gctgtaaagg	aaaaacacgc	c											21
<210>	6													
<211>	404													
<212>	DNA													
<213>	Homo sapiens													
<220>														
<221>	misc_feature													
<222>	(337)..(338)													
<223>	Non-consecutive bases													
<400>	6													
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	agggggcgaaa	tctctggaa	actggcagaa	ctcacagcca	atggcaggcg	ggagccgtcc								120

cgtagcgcc ggatccccgc gggtagggcg gggcgggcgg cgccgtgggg atcccggggc	180
agccgagggc ccctgactcg gtcctcgcg gcgacatgga tcggatggcc agctccatga	240
agcaggtgcc caacccactg cccaagggtgc tgagccggcg cggggtcggc gctgggctgg	300
aggcggcgga gcgcgagagc ttcgagcgga ctcaggttca gactgtcagc atcaataagg	360
ccattaatac gcagggaaagt ggctgtaaag gaaaaacatg ccag	404